

SEQUENCE LISTING

<110> Altosaar, Illimar
 Sardana, Ravinder
 Dudani, Aail
 Ganz, Peter
 Tackaberry, Eilleen

<120> Production of GM-CSF in Plants

<130> 08-898901US

<140> Not Yet Known

<141> 2003-11-26

<150> Canada 2,410,702

<151> 2002-11-26

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<170> PatentIn version 3.1

<210> 1

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<212> DNA

<213> Homo sapiens

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atg	gcg	cca	gca	ccg	agg	ccg	tcc	acc	cag	ccg	tgg	gag	cac	99
Met	Ala	Pro	Ala	Arg	Ser	Pro	Ser	Pro	Ser	Thr	Gln	Pro	Trp	His
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gtg	aac	gca	atc	cag	gag	gcc	cgc	agg	ctc	ctc	acc	ctc	tcc	cgc	gac	147
Val	Asn	Ala	Ile	Gln	Ala	Arg	Arg	Leu	Leu	Asn	Leu	Ser	Arg	Asp		
								35	40			45				

acc	gcc	gcc	gag	atg	aac	gag	acc	gtg	gag	gtg	atc	tcc	gag	atg	ttc	195
Thr	Ala	Ala	Glu	Met	Asn	Glu	Thr	Val	Glu	Val	Ile	Ser	Glu	Met	Phe	
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gat	ctc	cag	gag	ccg	acc	tgc	ctc	cag	acc	cgc	ctc	gag	ctg	tac	aag	243
Asp	Leu	Gln	Glu	Pro	Thr	Cys	Leu	Gln	Thr	Arg	Leu	Glu	Leu	Tyr	Lys	
						65		70		75						

cag	ggc	ctc	cgc	ggc	agc	ctc	acc	aag	ctc	aag	ggc	ccg	ctc	acc	atg	291
Gln	Gly	Leu	Arg	Gly	Ser	Leu	Thr	Lys	Leu	Lys	Gly	Pro	Leu	Thr	Met	
						80		85		90						

atg	gcg	tcc	cac	tac	aag	cag	cac	tgc	cca	ccg	acc	ccg	gag	acc	tcc	339
Met	Ala	Ser	His	Tyr	Lys	Gln	His	Cys	Pro	Pro	Thr	Pro	Glu	Thr	Ser	
95						100			105		110					

tgc gcc acc cag atc atc acc ttc gag agc ttc aag gag aac ctc aag	387
Cys Ala Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu Asn Leu Lys	
115 120 125	
gac ttc ctc ctc gtg atc ccg ttc gac tgc tgg gag ccg gtg cag gag	435
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35 40 45	
Ala Glu Met Asn Glu Thr Val Glu Val Ile Ser Glu Met Phe Asp Leu	
50 55 60	
Gln Glu Pro Thr Cys Leu Gln Thr Arg Leu Glu Leu Tyr Lys Gln Gly	
65 70 75 80	
Leu Arg Gly Ser Leu Thr Lys Leu Lys Gly Pro Leu Thr Met Met Ala	
85 90 95	
Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro Glu Thr Ser Cys Ala	
100 105 110	
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Pro Ala Arg Ser Pro Ser Pro Ser Thr Gln Pro Trp Glu His Val Asn
20           25           30

gcc atc cag gag gcc cgg cgt ctc ctg aac ctg agt aga gac act gct      144
Ala Ile Gln Ala Arg Arg Leu Leu Asn Leu Ser Arg Asp Thr Ala
35           40           45

gct gag atg aat gaa aca gta gaa gtg ata tca gaa atg ttt gac ctc      192
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50           55           60

cag gag ccc act tgc cta cag acc cgc ctg gag ctg tac aag cag ggc      240
Gln Glu Pro Thr Cys Leu Gln Thr Arg Leu Glu Leu Tyr Lys Gln Gly
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ctg cgg ggc agc ctc acc aag ctc aag ggc ccc ttg acc atg atg gcc      288
Leu Arg Gly Ser Leu Thr Lys Leu Lys Gly Pro Leu Thr Met Met Ala
85           90           95

agc cac tac aag cag cac tgc cct cca acc ccg gaa act tcc tgt gca      336
Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro Glu Thr Ser Cys Ala
100          105          110

acc cag att atc acc ttt gaa agt ttc aaa gag aac ctg aag gac ttc      384
Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu Asn Leu Lys Asp Phe
115          120          125

ctg ctt gtc atc ccc ttt gac tgc tgg gag cca gtc cag gag tga      429
Leu Leu Val Ile Pro Phe Asp Cys Trp Glu Pro Val Gln Glu
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20           25           30

Ala Ile Gln Glu Ala Arg Arg Leu Leu Asn Leu Ser Arg Asp Thr Ala

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Leu Arg Gly Ser Leu Thr Lys Leu Lys Gly Pro Leu Thr Met Met Ala
85 90 95

Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro Glu Thr Ser Cys Ala
100 105 110

Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu Asn Leu Lys Asp Phe
115 120 125

Leu Leu Val Ile Pro Phe Asp Cys Trp Glu Pro Val Gln Glu
130 135 140